

SEQUENCE LISTING

<110> Harris, Robert B.

Shen, Jane M.

Shahbaz, Manouchehr M.

<120> High Level Insect Expression of RAGE Proteins

<130> 41305-270555

<150> US 60/273,418

<151> 2001-03-05

<160> 10

<170> PatentIn version 3.1

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<212> DNA

<213> Homo sapiens

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aacgggtccc tcttccttc ggctgtcggg atccaggatg aggggatttt ccggtgccag      300
gcaatgaaca ggaatggaaa ggagaccaag tccaactacc gagtccgtgt ctaccagatt      360
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Trp	Gly	Ala	Val	Val	Gly	Ala	Gln	Asn	Ile	Thr	Ala	Arg	Ile	Gly	Glu		
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cca	ctg	gtg	ctg	aag	tgt	aag	ggg	gcc	ccc	aag	aaa	cca	ccc	cag	cgg		144
Pro	Leu	Val	Leu	Lys	Cys	Lys	Gly	Ala	Pro	Lys	Lys	Pro	Pro	Gln	Arg		
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Leu	Glu	Trp	Lys	Leu	Asn	Thr	Gly	Arg	Thr	Glu	Ala	Trp	Lys	Val	Leu		
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Gly	Lys	Pro	Leu	Val	Pro	Asn	Glu	Lys	Gly	Val	Ser	Val	Lys	Glu	Gln		
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acc	agg	aga	cac	cct	gag	aca	ggg	ctc	ttc	aca	ctg	cag	tcg	gag	cta		576
Thr	Arg	Arg	His	Pro	Glu	Thr	Gly	Leu	Phe	Thr	Leu	Gln	Ser	Glu	Leu		
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Met	Val	Thr	Pro	Ala	Arg	Gly	Asp	Pro	Arg	Pro	Thr	Phe	Ser	Cys			
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Ser	Phe	Ser	Pro	Gly	Leu	Pro	Arg	His	Arg	Ala	Leu	Arg	Thr	Ala	Pro		
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Ile	Gln	Pro	Arg	Val	Trp	Glu	Pro	Val	Pro	Leu	Glu	Glu	Val	Gln	Leu		
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ctg acc tgt gaa gtc cct gcc cag ccc tct cct caa atc cac tgg atg	816
Leu Thr Cys Glu Val Pro Ala Gln Pro Ser Pro Gln Ile His Trp Met	
260 265 270	
aag gat ggt gtg ccc ttg ccc ctt ccc ccc agc cct gtg ctg atc ctc	864
Lys Asp Gly Val Pro Leu Pro Leu Pro Pro Ser Pro Val Leu Ile Leu	
275 280 285	
cct gag ata ggg cct cag gac cag gga acc tac agc tgt gtg gcc acc	912
Pro Glu Ile Gly Pro Gln Asp Gln Gly Thr Tyr Ser Cys Val Ala Thr	
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cat tcc agc cac ggg ccc cag gaa agc cgt gct gtc agc atc agc atc	960
His Ser Ser His Gly Pro Gln Glu Ser Arg Ala Val Ser Ile Ser Ile	
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Leu Glu Trp Lys Leu Asn Thr Gly Arg Thr Glu Ala Trp Lys Val Leu
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Ser Pro Gln Gly Gly Gly Pro Trp Asp Ser Val Ala Arg Val Leu Pro
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Asn Gly Ser Leu Phe Leu Pro Ala Val Gly Ile Gln Asp Glu Gly Ile
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Phe Arg Cys Gln Ala Met Asn Arg Asn Gly Lys Glu Thr Lys Ser Asn
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Tyr Arg Val Arg Val Tyr Gln Ile Pro Gly Lys Pro Glu Ile Val Asp
115 120 125

Ser Ala Ser Glu Leu Thr Ala Gly Val Pro Asn Lys Val Gly Thr Cys
130 135 140

Val Ser Glu Gly Ser Tyr Pro Ala Gly Thr Leu Ser Trp His Leu Asp
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Gly Lys Pro Leu Val Pro Asn Glu Lys Gly Val Ser Val Lys Glu Gln
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Thr Arg Arg His Pro Glu Thr Gly Leu Phe Thr Leu Gln Ser Glu Leu
180 185 190

Met Val Thr Pro Ala Arg Gly Gly Asp Pro Arg Pro Thr Phe Ser Cys
195 200 205

Ser Phe Ser Pro Gly Leu Pro Arg His Arg Ala Leu Arg Thr Ala Pro
210 215 220

Ile Gln Pro Arg Val Trp Glu Pro Val Pro Leu Glu Glu Val Gln Leu
225 230 235 240

Val Val Glu Pro Glu Gly Gly Ala Val Ala Pro Gly Gly Thr Val Thr
245 250 255

Leu Thr Cys Glu Val Pro Ala Gln Pro Ser Pro Gln Ile His Trp Met
260 265 270

Lys Asp Gly Val Pro Leu Pro Leu Pro Pro Ser Pro Val Leu Ile Leu
275 280 285

Pro Glu Ile Gly Pro Gln Asp Gln Gly Thr Tyr Ser Cys Val Ala Thr
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